- 1 -

## SEQUENCE LISTING

(1)	GENERAL	INFORMATION:
<b>\</b>	(i)	APPLICANTS: SUZANNE L. TOPALIAN; STEVEN A. ROSENBERG; PAUL F. ROBBINS
	(ii)	TITLE OF INVENTION: MHC-CLASS II RESTRICTED MELANOMA ANTIGENS AND THEIR USE IN THERAPEUTIC METHODS
	(iii)	NUMBER OF SEQUENCES: 38
	(iv)	CORRESPONDENCE ADDRESS:  (A) ADDRESSEE: MORGAN & FINNEGAN, L.L.P.  (B) STREET: 345 PARK AVENUE  (C) CITY: NEW YORK  (D) STATE: NEW YORK  (E) COUNTRY: USA  (F) ZIP: 10154
	(v)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: FLOPPY DISK (B) COMPUTER: IBM PC COMPATIBLE (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: ASCII
	(vi)	CURRENT APPLICATION DATA:  (A) APPLICATION NUMBER: 08/533,895  (B) FILING DATE: 26 SEPT 1995  (C) CLASSIFICATION:
		ATTORNEY/AGENT INFORMATION: (A) NAME: FEILER, WILLIAM S. (B) REGISTRATION NUMBER: 26,728 (C) REFERENCE/DOCKET NUMBER: 2026-4205
	(ix)	TELECOMMUNICATION INFORMATION: (A) TELEPHONE: (212) 758-4800 (B) TELEFAX: (212) 751-6849 (C) TELEX: 421792
(2)	INFORMAT	TION FOR SEQ ID NO: :
	(i)	SEQUENCE CHARACTERISTICS:

- - (A) LENGTH: 15

  - (B) (C) (D) TYPE: AMINO ACID STRANDEDNESS: UNKNOWN
  - TOPOLOGY: UNKNOWN

	(xi)		SEQUENCE DESCRIPTION:						SEQ	ID NO:1		
Gln 1		Ile	Leu	Leu 5	Ser	Asn	Ala	Pro	Leu 10	Gly	Pro	
Gln	Phe	Pro 15										

- (2) INFORMATION FOR SEQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 14
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Asn Ile Leu Leu Ser Asn Ala Pro Leu Gly Pro Gln
1 5 10

Phe Pro

- (2) INFORMATION FOR SEQ ID NO:3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Gln Asn Ile Leu Leu Ser Asn Ala Pro Val Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

B1

Gln Asn Ile Leu Leu Ser Asn Val Pro Val Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Gln Asn Ile Leu Leu Ser Asn Val Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:6:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Asp Tyr Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

BI

Asp Tyr Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

Ser Gln Asp

- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Asp Gln Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp 15

- (2) INFORMATION FOR SEQ ID NO:9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Asp Phe Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:10:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Asp Tyr Ser Phe Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

274854\_1

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:11:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Asp Tyr Ser Tyr Leu Gln Asp Ser Val Pro Asp Ser 1 10

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:12:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 13
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Ser Tyr Leu Gln Asp Ser Val Pro Asp Ser Phe Gln
1 1 10

Asp

- (2) INFORMATION FOR SEQ ID NO:13:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 14
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Tyr Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser Phe 1 5 10

274854\_1

B1

## Gln Asp

- (2) INFORMATION FOR SEQ ID NO:14:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 13
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Ser Tyr Leu Gln Asp Ser Asp Pro Asp Ser Phe Gln 1 5 10

Asp

- (2) INFORMATION FOR SEQ ID NO:15:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Xaa Leu Leu Xaa Asn Xaa Xaa Leu Xaa

- (2) INFORMATION FOR SEQ ID NO:16:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 9
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Xaa Leu Gln Xaa Ser Xaa Xaa Asp Xaa 1 5

- (2) INFORMATION FOR SEQ ID NO:17:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN



	(xi)		SEQUENCE			ESCR:	ON:	SEQ	ID	NO:17		
Gln 1	Asn	Gln	Leu	Leu 5	Ser	Asn	Ala	Pro	Leu 10		У	Pro
Gln	Phe	Pro 15										

- (2) INFORMATION FOR SEQ ID NO:18:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Gln Asn Phe Leu Leu Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:19:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Gln Asn Val Leu Leu Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:20:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN



(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Gln Asn Ile Gln Leu Ser Asn Ala Pro Leu Gly Pro

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:21:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Gln Asn Ile Phe Leu Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:22:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Gln Asn Ile Val Leu Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:23:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

B

Gln Asn Ile Leu Gln Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:24:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Gln Asn Ile Leu Phe Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:25:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Gln Asn Ile Leu Val Ser Asn Ala Pro Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:26:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Gln Asn Ile Leu Leu Ser Asn Gln Pro Leu Gly Pro 1 5 10

274854\_1

BI

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:27:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Gln Asn Ile Leu Leu Ser Asn Ala Gln Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:28:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Gln Asn Ile Leu Leu Ser Asn Ala Val Leu Gly Pro 1 5 10

Gln Phe Pro 15

- (2) INFORMATION FOR SEQ ID NO:29:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

Gln Asn Ile Leu Leu Ser Asn Ala Pro Gln Gly Pro 1 5 10

274854\_1

- (C) STRANDEDNESS: UNKNOWN
  (D) TOPOLOGY: UNKNOWN

  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

  Ser Asn Ala Pro Leu Gly Pro Gln Phe Pro Phe
  1 5 10

  Thr Gly Val Asp
  15

  (2) INFORMATION FOR SEQ ID NO:34:

  (i) SEQUENCE CHARACTERISTICS:
  - (B) TYPE: AMINO ACID (C) STRANDEDNESS: UNKNOWN

LENGTH: 15

- (D) TOPOLOGY: UNKNOWN
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

Asp Tyr Ser Gln Leu Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp

(2) INFORMATION FOR SEQ ID NO:35:

(A)

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15
  - (B) TYPE: AMINO ACID
  - (C) STRANDEDNESS: UNKNOWN
  - (D) TOPOLOGY: UNKNOWN
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

Asp Tyr Ser Tyr Gln Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:36:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN

BI

- (D) TOPOLOGY: UNKNOWN
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

Asp Tyr Ser Tyr Phe Gln Asp Ser Asp Pro Asp Ser 1 5 10

Phe Gln Asp

- (2) INFORMATION FOR SEQ ID NO:37:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 12
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

Tyr Leu Gln Asp Ser Asp Pro Asp Ser Phe Gln Asp
1 5 10

- (2) INFORMATION FOR SEQ ID NO:38:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 11
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS: UNKNOWN
    - (D) TOPOLOGY: UNKNOWN
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Leu Gln Asp Ser Asp Pro Asp Ser Phe Gln Asp
1 5 10





Creation date: 11-17-2003

Indexing Officer: RYESHITILA - RAHEL YESHITILA

Team: OIPEBackFileIndexing

Dossier: 08533895

Legal Date: 06-22-2000

No.	Doccode	Number of pages
1	NPL	9

Total number of pages: 9
Remarks:
Order of re-scan issued on